

REMARKS

Claims 1-17, 19-32, 52-57, 59, 73, and 76-92 are currently pending in the present application.

Applicants note that the Examiner has withdrawn objections to the drawings and specification and to claims 3 and 30 due to minor informalities or typographical errors. The rejection of claims 5-6, 8, 11-12, and 26-27 under 35 U.S.C. §112, second paragraph has also been withdrawn.

Claim 1 has been amended to recite “the recombinant protein variant” after the second occurrence of “naturally occurring allergen.” Support for this amendment may be found throughout the specification and particularly in original claims 1, and 52-54.

Claims 25-26 and 30-32 have been amended to include reference to Accession Numbers. Support for these amendments may be found in the specification on at least pages 30 and 32.

Claims 52-54 have been amended to recite “comprises two or more primary mutations spaced by at least one non-mutated amino acid residue.” Support for this amendment may be found throughout the specification and in original claims 1 and 19.

No new matter has been introduced in these amendments. Entry and consideration of these amendments is respectfully requested.

Amendments To The Specification

The specification has been amended at page 30, line 16; and at page 31, line 13 to replace “Q76H” with “E76Q.” Support for this amendment may be found in the specification on page 30 and in Fig. 21 where residue 76 is described as being glutamic acid (E) in the reference sequence Accession No. AJ488060.

The Rejections Under the Second Paragraph of 35 U.S.C. § 112 Should Be Withdrawn

Claims 1-17, 19-32, 52-57, 59, 73, and 76-92 have been rejected under 35 U.S.C. § 112, second paragraph, as being indefinite. In particular, the Office Action indicates that claim 1 is

rejected claims 54 and 73 under 35 U.S.C. § 102(b) as anticipated by the publication of King *et al.*, J. Immun., 2001, 166(10):6057-6065 (“King”).

A. The Legal Standard of Anticipation

Anticipation requires that each and every element of the rejected claim(s) be disclosed in a single prior art reference. See M.P.E.P. §2131 (8th Ed. Rev. 2, May 2004). “A claim is anticipated only if each and every element as set forth in the claim is found, either expressly or inherently described, in a single prior art reference.” *Verdegaal Bros. v. Union Oil Co. of California*, 814 F.2d 628, 631, 2 USPQ2d 1051, 1053 (Fed. Cir. 1987). Every element of the claimed invention must literally present, arranged as in the claim. *Perkin Elmer Corp. v. Computervision Corp.*, 732 F.2d 888, 894, 221 USPQ 669, 673 (Fed. Cir. 1984).

B. Valenta Does Not Anticipate the Pending Claims

Valenta does not teach the use of a modified scaffold protein with a similar three-dimensional folding pattern to that of a desired natural allergen, nor does Valenta teach the insertion of mutations in the scaffold protein as recited in claim 1. Valenta teaches recombinant allergen proteins. In contrast, the present invention teaches the use of a scaffold protein which maintains the three-dimensional folding pattern of the allergen, and the introduction of point mutations into the scaffold protein, not into the allergen itself. Applicants note that the present claims do not encompass recombinant scaffold proteins that are identical to the allergen. Instead, the inventive recombinant scaffold proteins are modified when compared to the unmodified scaffold protein (i.e. the template) as well as when compared to the naturally occurring allergen.

Additionally, the claims require that the recombinant protein exhibit increased affinity and/or binding capacity to IgE antibodies specific to the naturally occurring allergen. Valenta does not teach recombinant mutant proteins that meet this requirement. Valenta does not teach recombinant proteins with structural similarity, *i.e.*, proteins with a similar tertiary structure. Applicants note that the Valenta results from IgE immunoblots are indicative only of binding conferred by an antigen binding site. See Valenta, col. 3, line 56 to col. 4, line 9. Having similar binding may or may not be due to a protein having an overall similar 3-D structure; it may be conferred by a protein with an antigenic site that confers this binding, while the protein does not

